Amendments to the claims:

This listing of claims is to replace all prior versions and listings of claims in the application.

- 1. (Withdrawn) An isolated nucleic acid molecule comprising a sequence substantially identical to SEQ ID NO:252.
 - 2-43. (Cancelled).
- 44. (Withdrawn) A substantially pure polypeptide comprising an amino acid sequence that is substantially identical to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 92 (ORF11738).
- 45. (Withdrawn) The polypeptide of claim 44, comprising an amino acid sequence that is identical to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 92.
- 46. (Withdrawn) A substantially pure polypeptide comprising an amino acid sequence that is substantially identical to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 220 (ORF14155).

- 47. (Withdrawn) The polypeptide of claim 46, comprising an amino acid sequence that is identical to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 220.
- 48. (Currently Amended) A substantially pure polypeptide comprising an amino acid sequence that has a least 90% identity has at least 90% identity to the amino acid sequence of a polypeptide the full-length polypeptide encoded by SEQ ID NO: 252, wherein said polypeptide is a virulence factor.
- 49. (Original) The polypeptide of claim 48, comprising an amino acid sequence that is identical to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 252.
- 50. (Withdrawn) A substantially pure polypeptide comprising an amino acid sequence that is substantially identical to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 272 (ORF23228).
- 51. (Withdrawn) The polypeptide of claim 50, comprising an amino acid sequence that is identical to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 272.

- 52. (Withdrawn) A method for identifying a compound which binds a polypeptide, said method comprising the steps of:
- (a) contacting a candidate compound with a substantially pure polypeptide of claim 44, 46, 48, or 50 under conditions that allow binding; and
 - (b) detecting binding of the candidate compound to the polypeptide.
- 53. (Previously presented) The substantially pure polypeptide of claim 48, wherein said polypeptide has a least 95% identity to the amino acid sequence of a polypeptide encoded by SEQ ID NO: 252.